

GenCore version 5.1.6  
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OM protein - protein search, using sw model  
Run on: March 7, 2005, 06:55:26 ; Search time 46.7139 Seconds  
(without alignments)  
919.008 Million cell updates/sec

Title: US-09-939-537-37  
Perfect score: 591  
Sequence: 1 TRFSRSARPPAYQOGQNDQY.....LSTATKDYDALHMQALPPR 111  
Scoring table: BLOSUM62  
Gapop 10.0 , Gapext: 0.5

Searched: 2105692 seqs, 386761381 residues

Total number of hits satisfying chosen parameters: 2105692  
Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : A\_Geneseq\_16Dec04:  
1: geneseqP1990s: \*  
2: geneseqP2000s: \*  
3: geneseqP2001s: \*  
4: geneseqP2002s: \*  
5: geneseqP2003as: \*  
6: geneseqP2003bs: \*  
7: geneseqP2004s: \*  
8: geneseqP2004bs: \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match Length	DB ID	Description
1	591	100.0	111	AAR78675
2	591	100.0	111	AAR89452
3	583	98.6	142	AAR0226
4	583	98.6	142	AAR14196
5	583	98.6	163	AAR76486
6	583	98.6	443	ABP76486
7	583	98.6	443	ABP74240
8	583	98.6	532	AAR27278
9	583	98.6	532	AAR78678
10	583	98.6	532	AAR89458
11	583	98.6	532	AAR02215
12	583	98.6	2	AAR8311
13	572	96.8	112	ADL67239
14	572	96.8	8	ADL67237
15	572	96.8	163	ABO84748
16	572	96.8	357	ABO73652
17	572	96.8	395	ABG73653
18	572	96.8	444	AAR36845
19	572	96.8	473	AAR26646
20	572	96.8	514	AAR26647
21	572	96.8	512	AAW73051
22	572	96.8	631	ADL34696
23	572	96.8	623	AAR84965
24	572	96.8	634	ABP82300
25	572	96.8	643	AAW73050

Raw73048 A33 chime  
Raw26649 Chimeric  
Raw26650 Chimeric  
Ab084747 Human can  
Aay52340 Full-leng  
Abp08920 Mouse tru  
Adh74813 Zeta. 4/2  
Adh74805 Mouse imm  
Abp84745 Mouse can  
Aar85508 Leader-sc  
Raw82315 Chimeric  
Aay52345 Mutant fu  
Abp08925 Mouse tru  
Adh74810 CD5/Zeta  
Raw24027 Single ch  
Raw24025 Single ch  
Aay52344 Mutant fu  
Abp08924 Mouse tru  
Adh74809 CDB/Zeta  
Raw13273 Murine CD

#### ALIGNMENTS

RESULT 1
AAR78675
ID AAR78675 standard; protein, 111 AA.
XX
AC AAR78675;
XX
DT 12-APR-1996 (first entry)
DB T-cell receptor Protein zeta intracellular domain.
XX
KW Chimeric receptor; CD4; T-cell receptor zeta; HIV; cytolysis; human immunodeficiency virus; adoptive immunotherapy.
XX
OS Homo sapiens.
XX
PN W09521528-A1.
PD
XX
17-AUG-1995.
PR 12-JAN-1995;
XX
PR 14-FEB-1994;
XX
PR 02-AUG-1994;
XX
(GEHO ) GEN HOSPITAL CORP.
PT Seed B, Banapour B, Romeo C, Kolanus W;
XX
DR WI: 1995-292893/38.
DR N-PSDB; AAQ6105.
XX
PT Target cytolytic of HIV-infected cells - by chimeric CD4 receptor-bearing cells.
XX
PS Example 10; Fig 27; 118pp; English.
XX
The intracellular domain (AAR78675) of human T-cell receptor protein zeta is used in the construction of a chimeric receptor utilised in the targeted cytolysis of cells expressing HIV envelope proteins on their surface. The chimeric receptor comprises the extracellular domain (pref. amino acids 1-394 or 1-200) of CD4 linked to the intracellular portion of e.g. zeta
XX
Sequence 111 AA;

Query Match Similarity 100.0%; Score 591; DB 2; Length 111; Best Local Similarity 100.0%; Pred. No. 1.6e-59; Matches 111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	1 TRFSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 60	QY	61 LOKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 111
Db	1 TRFSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 60	Db	61 LOKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 111
QY	61 LOKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 111	RESULT 3	
Db	61 LOKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 111	AC	AAR89461;
RESULT 2		XX	
ID	AAR89452 standard; peptide; 111 AA.	DT	26-SEP-1996 (first entry)
XX		XX	
AC	AAR89452;	DE	zeta intracellular domain.
XX		XX	
DT	26-SEP-1996 (first entry)	KW	CD7; transmembrane domain; chimeric receptor; CD5; CD34; CH2; CH3; IgG1; human; CD4; HIV; proteinaceous alpha-helix; T cell; B cell; neutrophil; dendritic cell; therapy; mammal; infection.
XX		KW	
DB		OS	Homo sapiens.
XX		OS	
KW		XX	
human; CD4; HIV; proteinaceous alpha-helix; T cell; B cell; neutrophil; dendritic cell; therapy; mammal; infection.		PN	W09603883-A1.
XX		XX	
XX		PD	15-FEB-1996.
XX		PP	26-JUL-1995; 95WO-US009468.
PN	W09603883-A1.	XX	
XX		PR	02-AUG-1994; 94US-00284391.
XX		PR	24-FEB-1995; 95US-00394388.
PD	15-FEB-1996.	XX	
PP	26-JUL-1995; 95WO-US009468.	PA	(GEHO ) GEN HOSPITAL CORP.
XX		XX	
PR	02-AUG-1994; 94US-00284391.	PA	Seed B, Banapour B, Romeo C, Kolanus W;
PR	24-FEB-1995; 95US-00394388.	PI	
XX		XX	
PA	(GEHO ) GEN HOSPITAL CORP.	DR	WPI; 1996-129034/13.
XX		XX	
PT	Seed B, Banapour B, Romeo C, Kolanus W;	PT	Membrane-bound chimeric receptor comprising extracellular portion including CD4 fragment - cells expressing receptor can be used for treatment of HIV infection.
XX		PT	
DR	WPI; 1996-129034/13.	XX	
N-PSDB; AAT10799.		PS	Example 8; Page 96; 134pp; English.
XX		XX	
PT		CC	This sequence represents the zeta intracellular domain. This sequence is included in the membrane bound proteinaceous chimeric receptor of the invention. The extracellular portion of the chimeric receptor contains a fragment of CD4 (amino acids 1-394 or 1-200 of the CD4 sequence see AAR89450 and AAR89451) which specifically recognises and binds HIV-infected cells, but does not mediate HIV infection. The extracellular domain of the receptor is separated from the cell membrane by 48 or 72 angstroms, or by one or more proteinaceous alpha-helices. The transmembrane region of the chimeric receptor contains a portion of the CD7 (see AAR89440), CD5 or CD34 transmembrane domain. Alternatively, the extracellular portion of the receptor can also be separated from the intracellular domain by the hinge, CH2 and CH3 domains of human IgG1 (see AAR89441). The cells expressing the receptor are preferably T cells, B cells, neutrophils, or dendritic cells. The therapeutic cells expressing the chimeric receptor are administered to a mammal to treat HIV infection
XX		CC	
PS	Example 10; Fig 27; 134pp; English.	XX	
XX		CC	This sequence represents the zeta intracellular domain. This sequence is included in the membrane bound proteinaceous chimeric receptor of the invention. The extracellular portion of the chimeric receptor contains a fragment of CD4 (amino acids 1-394 or 1-200 of the CD4 sequence see AAR89450 and AAR89451) which specifically recognises and binds HIV-infected cells, but does not mediate HIV infection. The extracellular domain of the receptor is separated from the cell membrane by 48 or 72 angstroms, or by one or more proteinaceous alpha-helices. The transmembrane region of the chimeric receptor contains a portion of the CD7 (see AAR89440), CD5 or CD34 transmembrane domain. Alternatively, the extracellular portion of the receptor can also be separated from the intracellular domain by the hinge, CH2 and CH3 domains of human IgG1 (see AAR89441). The cells expressing the receptor are preferably T cells, B cells, neutrophils, or dendritic cells. The therapeutic cells expressing the chimeric receptor are administered to a mammal to treat HIV infection
XX		CC	
PS	Sequence 111 AA;	XX	
XX		CC	Sequence 142 AA;
Query Match	100.0%; Score 591; DB 2; Length 111;	CC	Query Match 98.6%; Score 583; DB 2; length 142;
Best Local Similarity	100.0%; Pred. No. 1.6e-59; Mismatches 0; Indels 0; Gaps 0;	CC	Best Local Similarity 99.1%; Pred. No. 1.8e-58; Mismatches 0; Indels 0; Gaps 0;
Matches	111; Conservative	CC	Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
QY	1 TRFSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 60	QY	2 RRSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 61
Db	1 TRFSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 60	Db	33 KRSRSAEPAYQOCONQNLNELNLRREYDVLKRRGRDPENGKPRKRKNPOEGLYNE 92
QY	62 QDKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 111	QY	62 QDKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 142
Db	63 QDKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 142	Db	93 QDKKMAEAYSEIGMGERRGKGHDGLXQGLSTATOTYDALHMQALPPR 142

**RESULT 4**

ID AAW02226 Standard; peptide; 142 AA.

AC AAW02226;

XX XX

DT 11-NOV-1996 (first entry)

DE T-cell receptor zeta chain.

XX XX

KW Chimaeric receptor; cellular immunity; adoptive immunotherapy; CD16; human immunodeficiency virus type 1; HIV-1; AIDS; therapy; T-cell receptor zeta chain; cytotoxic T lymphocyte; CTL.

OS Homo sapiens.

PN W09625953-A1.

PD 29-AUG-1996.

PP 25-JAN-1995; 96WO-US001056.

PR 24-FEB-1995; 95US-00394176.

PA (GEHO ) GEN HOSPITAL CORP.

XX PA

PI Seed B, Romeo C, Kolanus W;

XX DR WPI; 1999-340518/29.

DR N-PSDB; AAXK1131.

XX PT New T cell receptor zeta chain protein - useful for detecting auto-immune disease.

XX RS Claim 3; Page 12-13; 22pp; Japanese.

XX CC This sequence is the T cell receptor zeta chain protein of the invention, which is used for detecting an autoimmune disease in a patient. The protein can be used for detecting autoimmune disease affliction. Systemic lupus erythematosus (SLE) can be diagnosed by detecting the mutation of T cell zeta chain gene and can be treated by complementing the detected mutation with a normal zeta chain gene or zeta chain protein

XX SQ Sequence 163 AA;

Query Match 98.6%; Score 583; DB 2; Length 163;

Best Local Similarity 99.1%; Pred. No. 2.1e-58; Mismatches 0; Indels 0; Gaps 0;

Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 2 RESRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 61

Db 54 KFRSRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 113

OY 62 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 111

Db 114 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 163

CC Sequence 142 AA;

XX SQ Sequence 142 AA;

Query Match 98.6%; Score 583; DB 2; Length 142;

Best Local Similarity 99.1%; Pred. No. 1.8e-58; Mismatches 0; Indels 0; Gaps 0;

Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 2 RESRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 61

Db 33 KFRSRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 92

OY 62 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 111

Db 93 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 142

RESULT 5

ID AY14196

AC AY14196 standard; protein; 163 AA.

XX XX

AC AY14196;

XX XX

DT 27-JUL-1999 (first entry)

XX XX

DE T cell receptor zeta chain protein sequence.

XX XX

KW T cell receptor zeta chain protein; autoimmune disease; diagnosis; SLE; systemic lupus erythematosus; therapy.

OS Homo sapiens.

PN JP1121082-A.

XX PD 11-MAY-1999.

XX PR 23-OCT-1997; 97JP-00309302.

XX PR 23-OCT-1997; 97JP-00309302.

XX PA (TAKE-) TAKEUCHI T.

PA (NOBE-) NOBEL IGAKU KENKYUSHO YG.

XX DR WPI; 1999-340518/29.

DR N-PSDB; AAXK1131.

XX PT New T cell receptor zeta chain protein - useful for detecting auto-immune disease.

XX RS Claim 3; Page 12-13; 22pp; Japanese.

XX CC This sequence is the T cell receptor zeta chain protein of the invention, which is used for detecting an autoimmune disease in a patient. The protein can be used for detecting autoimmune disease affliction. Systemic lupus erythematosus (SLE) can be diagnosed by detecting the mutation of T cell zeta chain gene and can be treated by complementing the detected mutation with a normal zeta chain gene or zeta chain protein

XX SQ Sequence 163 AA;

Query Match 98.6%; Score 583; DB 2; Length 163;

Best Local Similarity 99.1%; Pred. No. 2.1e-58; Mismatches 0; Indels 0; Gaps 0;

Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 2 RESRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 61

Db 54 KFRSRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 113

OY 62 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 111

Db 114 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 163

CC Sequence 142 AA;

XX SQ Sequence 142 AA;

Query Match 98.6%; Score 583; DB 2; Length 142;

Best Local Similarity 99.1%; Pred. No. 1.8e-58; Mismatches 0; Indels 0; Gaps 0;

Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 2 RESRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 61

Db 33 KFRSRSABEPAYQOGQONQLYNELNLRGRRREYDVLKDRGRDPENGKERRRKPQEGLYNEL 92

OY 62 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 111

Db 93 QDKKMAEAYSEIGMKGRRRGKDHGLYQGLSTATKOTYDALHMQALPPR 142

RESULT 6

ID ABG76488

AC ABG76488 standard; protein; 443 AA.

XX XX

AC ABG76488;

XX DT 13-MAY-2003 (first entry)

DB Humanised anti-CEA antibody, HMN14.

XX CEA: carcinoembryonic antigen; IgTcR; T-cell receptor; cancer; tumour; colorectal cancer; breast cancer; lung cancer; HMN14; cytostatic; mouse; human; zeta signalling chain; CD8alpha hinge; humanised antibody.

XX OS Homo sapiens.

OS Mus sp.

OS Synthetic.

OS Chimeric.

XX PN US2002165360-A1.

XX PD 07-NOV-2002.

XX PR 10-DEC-2001; 2001US-00006771.

XX PR 30-NOV-2000; 2000US-0250087P.

PR 30-NOV-2000; 2000US-0250090P.

PA	(JUNG/)	JUNGHANS R P.
XX	XX	PR 30-NOV-2000; 2000US-0250089P.
PX	XX	PT (JUNG/)
PI	XX	PT JUNGHANS R P.
XX	XX	PT
WPI:	2003-298705/29.	PT
DR	N-PSDB; ABX13168.	PT
XX	XX	PT New chimeric molecule from humanized antibody against carcinoembryonic antigen and having signaling molecules of T cells and other effector cells, useful for the treatment of colorectal, breast and lung cancers.
PS	PS Disclosure; Page 7-8; 20pp; English.	PT
XX	CC The invention relates to a chimaeric molecule comprising the carcinoembryonic antigen (CEA) binding domain of humanised antibody MN14 as a single chain antibody with a (GSGS) <sub>3</sub> linker, the zeta signalling chain of the T cell receptor (TCR) and an intervening CDBalpha hinge in which the cysteine residues have been mutated, with the IgTCR molecule occupying nucleotides 2426-3766 of the retroviral vector sequence. The new chimaeric molecule expressed in T cells, NK (not defined) or other effector cells are useful in treating patients with cancers expressing the CEA antigen, together with other or with heterologous constructs to engage additional stimulatory and functional properties of the effector cells to enhance the anti-tumour therapeutic efficacy. The cancer disorder includes colorectal, breast and lung cancers. The present sequence represents the chimaeric molecule of the invention	PT
XX	CC Sequence 443 AA;	PT
QY	Query Match 98.6%; Score 583; DB 6; Length 443; Best Local Similarity 99.1%; Pred. No. 7.3e-58; Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0; Sequence 443 AA;	PT
Db	334 KFRSRAEPPAYQQGQNQLYNELNIGREYVLDVLRGRDPMEGGKPRRNPOEGLYNEL 61	PT
QY	2 RFRSRAEPPAYQQGQNQLYNELNIGREYVLDVLRGRDPMEGGKPRRNPOEGLYNEL 61	PT
Db	62 QDKDMEAAYSEIGMKGERRGKGHDGLYQGLSTATKDTYDALHMQALPPR 111	PT
QY	394 QDKDMEAAYSEIGMKGERRGKGHDGLYQGLSTATKDTYDALHMQALPPR 443	PT
RESULT 7		PT
ABG74240		PT
ID	ABG74240 standard; protein; 443 AA.	PT
AC	ABG74240;	PT
XX	DT 23-OCT-2003 (revised)	PT
DT	22-APR-2003 (first entry)	PT
XX	DE ChimERIC HMM14/T-cell receptor.	PT
XX	DE Retroviral vector; T-cell receptor; HMM14; antibody; IgTCR; receptor; cytosolic; dermatological; neuroprotective; immunomodulant; Gd3; ganglioside antigen; MB3.6; PSMA; tumour; 3D8; 4D4; 3E11; prostate-specific membrane antigen; zeta; signalling chain; human; cancer; melanoma; neuroendocrine tumour; prostate cancer; small cell lung cancer; mouse; CD8alpha hinge.	PT
KW	OS Homo sapiens.	PT
OS	Mus sp.	PT
OS	Chimeric.	PT
PN	US2003132983-A1.	PT
XX	DD 19-SEP-2002.	PT
XX	XX	PT
XX	10-DEC-2001; 2001US-00006773.	PT
XX	30-NOV-2000; 2000US-0250087P.	PT
XX	XX	PR N-PSDB; ABX16565.
XX	XX	PT New chimeric molecule useful in treating patients with disorders, such as melanoma, neuroendocrine disorders, prostate and small cell lung cancer comprises GD3 and/or PSMA binding domains of antibody.
XX	PS Disclosure; Page 7-8; 35pp; English.	PT
XX	CC The invention relates to a chimaeric molecule comprising the GD3 (ganglioside antigen) binding domain of antibody MB3.6, with any of 3 variable gene sequences, or the PSMA (prostate-specific membrane antigen) binding domain of antibody 3D8, 4D4 and 3E11, with variable gene sequences, the zeta signalling chain of the T cell receptor and an intervening CDBalpha hinge in which cysteine residues have been mutated. The chimeric molecules expressed in T cells or NK cells or other effector cells are useful in treating patients with cancers expressing the GD3 (MB3.6 derivatives) or PSMA antigen (3D8, 4D4, 3E11 derivatives), and/or together with each other or with heterologous constructs to engage additional stimulatory and functional properties of the effector cells to enhance the antitumour therapeutic efficacy (claimed). They are particularly useful in disorders including melanoma, neuroendocrine tumours and prostate and small cell lung cancer. The present sequence is modified CDBalpha hinge and the T-cell receptor zeta chain (IgTCR) encoded by a retroviral vector. The hM14 antibody coding region is replaced with the MB3.6, 3D8, 4D4 or 3E11 genes of the invention. (Updated on 23-OCT-2003 to standardise OS field)	PT
XX	CC Sequence 443 AA;	PT
QY	Query Match 98.6%; Score 583; DB 6; Length 443; Best Local Similarity 99.1%; Pred. No. 7.3e-58; Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0; Sequence 443 AA;	PT
Db	334 KFRSRAEPPAYQQGQNQLYNELNIGREYVLDVLRGRDPMEGGKPRRNPOEGLYNEL 393	PT
QY	2 RFRSRAEPPAYQQGQNQLYNELNIGREYVLDVLRGRDPMEGGKPRRNPOEGLYNEL 61	PT
Db	62 QDKDMEAAYSEIGMKGERRGKGHDGLYQGLSTATKDTYDALHMQALPPR 111	PT
QY	394 QDKDMEAAYSEIGMKGERRGKGHDGLYQGLSTATKDTYDALHMQALPPR 443	PT
RESULT 8		PT
AAR27278		PT
ID	AAR27278 standard; protein; 512 AA.	PT
XX	AC AAR27278;	PT
XX	DT 25-MAR-2003 (revised)	PT
DT	28-JUL-1995 (first entry)	PT
XX	DE CD4:gamma peptide chimeric protein.	PT
XX	DE Fusion protein; CD4; extracellular domain; zeta; eta; gamma; membrane spanning domain; intracellular domain; type I; integral membrane homodimer; TCR; T cell antigen receptor; extracellular domain; mouse; human; receptor; chimera; HPB-ALL tumour cell line; natural killer cell.	PT
XX	OS Homo sapiens.	PT
PN	W09215322-A1.	PT
XX	DD 17-SEP-1992.	PT

XX 06-MAR-1992; 92WO-US001785.  
 PR 07-MAR-1991; 91US-00665961.  
 XX (GEHO ) GEN HOSPITAL CORP.  
 PA  
 XX Seed B, Romeo C, Kolanus W;  
 DR WPI; 1992-331474/40.  
 XX N-PSDB; AAQ28706.  
 PT Therapeutic cells expressing chimeric receptors - directing cellular response to an infective agent, useful in treating HIV-1, AIDS  
 XX Pneumocystis carinii infections etc.  
 PS Example 2; Page 74-76; 114pp; English.  
 XX This sequence represents a fusion protein between the CD4 extracellular domain and the gamma protein membrane spanning domain and intracellular domain. The Fc-receptor-associated gamma chain is expressed in cell surface complexes with additional polypeptides, some of which mediate ligand recognition, and others which have undefined function. Gamma bears a homodimeric structure and overall organisation very similar to that of zeta (see also A028704), and is a component of both the mast cell/basophil high affinity IgE receptor, Fc-epsilon RI, which consists of at least three distinct polypeptide chains and one of the low affinity receptors for IgG, represented in mice by Fc-gamma-RII-alpha. In the production of the CD4 receptor chimera, the gamma cDNA was isolated from the HPB-ALL tumour cell line and from human natural killer cells. The gamma cDNA was joined to the extracellular domain by engineering a BamHI site naturally present a few residues upstream of the membrane spanning domain. (Updated on 25-MAR-2003 to correct PN field.)  
 CC Sequence 532 AA;  
 SQ  
 Query Match 98.6%; Score 583; DB 2; Length 532;  
 Best Local Similarity 99.1%; Pred. No. 9.2e-58; Indels 0; Gaps 0;  
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 AC AAR89458;  
 XX AAR89458 standard; protein; 532 AA.  
 ID AAR78678  
 XX AAR78678;  
 AC AAR78678;  
 DT 16-APR-1996 (first entry)  
 KW T-cell receptor eta.  
 KW Chimeric receptor; CD4; T-cell receptor eta; HIV; cytolysis; human immunodeficiency virus; adoptive immunotherapy.  
 OS Homo sapiens.  
 XX WO9521528-A1.  
 PN  
 PR 12-JAN-1995; 95WO-US00454.  
 PR 14-FEB-1994; 94US-00195395.  
 PR 02-AUG-1994; 94US-00284391.  
 XX  
 PT Target cytolytic of HIV-infected cells - by chimeric CD4 receptor-bearing cells.  
 XX N-PSDB; AAQ6124.  
 PT Fusion proteins comprising the extracellular domain of CD4 fused to T-cell receptor zeta, gamma or eta (A028706-78, respectively) were expressed in CV1 using a vaccine virus vector. These CD4:zeta, CD4:gamma and CD4:eta chimeric receptors mediated cytolytic of targets expressing HIV gp120/41.  
 CC Sequence 532 AA;  
 SQ  
 Query Match 98.6%; Score 583; DB 2; Length 532;  
 Best Local Similarity 99.1%; Pred. No. 9.2e-58; Indels 0; Gaps 0;  
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 AC AAR89458;  
 XX AAR89458 standard; protein; 532 AA.  
 ID AAR78678  
 XX AAR78678;  
 AC AAR78678;  
 DT 26-SEP-1996 (first entry)  
 DE CD4:eta fusion protein.  
 KW CD7; transmembrane domain; chimeric receptor; CD5; CD4; CD2; CT1; IgG1; KW human; CD4; HIV; proteinaceous; alpha-helix; T cell; B cell; neutrophil; KW dendritic cell; therapy; mammal; infection.  
 XX OS Synthetic.  
 XX PN WO9603883-A1.  
 XX PD 15-FBB-1996.  
 XX PR 26-JUL-1995; 95WO-US009468.  
 XX PR 02-AUG-1994; 94US-00284391.  
 XX PR 24-FEB-1995; 94US-00394388.  
 PA (GEHO ) GEN HOSPITAL CORP.  
 XX PI Seed B, Banapour B, Romeo C, Kolanus W;  
 XX DR WPI; 1996-129034/13.  
 XX DR N-PSDB; A0110803.  
 PT Membrane-bound chimeric receptor comprising extracellular portion including CD4 fragment - cells expressing receptor can be used for treatment of HIV infection.  
 XX  
 PS Example 2; Page 80-81; 134pp; English.  
 XX AAT0801-T10803 represent membrane bound proteinaceous chimeric receptors of the invention. This sequence represents the CD4:eta chimera. The  
 CC

CC transmembrane region of the chimeric receptor acts to separate the  
 intracellular and extracellular domains of the chimera, and contains a  
 portion of the CD7 (see AAR8940), CD5 or CD34 transmembrane domains.  
 Alternatively, the extracellular portion of the receptor can be separated  
 from the intracellular domain by the hinge, CH2 and CH3 domains of human  
 IgG1 (see AAR9441). The extracellular portion of the chimeric receptor  
 contains a fragment of CD4 (amino acids 1-394 or 1-200 of the CD4  
 sequence, see AAR89450 and AAR9451) which specifically recognises and  
 binds HIV-infected cells, but does not mediate HIV infection. The  
 extracellular domain of the receptor is separated from the cell membrane  
 by 48 or 72 angstroms, or by one or more proteinaceous alpha-helices. The  
 cells expressing the receptor are preferably T cells, B cells,  
 neutrophils, or dendritic cells. The therapeutic cells expressing the  
 chimeric receptor are administered to a mammal to treat HIV infection  
 XX Sequence 532 AA;  
 SQ

Query	Match	Best Local Similarity	Score	DB	Length
Qy	2 RFSRSRAPPYQQGSQNLQYKELNLRGRREYVLDKRGDRDEMGKRRRNQPEGLYNEL	98 %	583	2;	532;
Db	423 KPSRSRAPPYQQGSQNLQYKELNLRGRREYVLDKRGDRDEMGKRRRNQPEGLYNEL	99.1%	1;	Mismatches	0;
Qy	62 QDKDRAEAVSEIGMKGERRRKGHGHLIQGUSTATKOTYDALHMQALPPR	98 %	111		
Db	483 QDKDRAEAVSEIGMKGERRRKGHGHLIQGUSTATKOTYDALHMQALPPR	98 %	532		

**RESULT 11**

ID	AAW02215 standard; protein; 532 AA.
AC	AAW02215;
DT	16-OCT-2003 (revised) 11-NOV-1996 (first entry)
DE	CD4: T-cell receptor eta chain chimeric receptor.
XX	
KW	chimeric receptor; cellular immunity; adoptive immunotherapy; CD4; human immunodeficiency virus type 1; HIV-1; AIDS; therapy; T-cell receptor eta chain; cytotoxic T lymphocyte; CTL.
XX	
OS	Homo; sapiens.
OS	Mus sp.
OS	Chimeric.
XX	
FT	Key
FT	Domain
FT	Region
FT	Region
FT	Region
FT	Location/Qualifiers
FT	1. .393
FT	/label= Extracellular domain
FT	/note= "CD4 extracellular domain"
FT	394. .396
FT	/label= Linker
FT	/note= "encoding DNA contains a BamHI site used for fusion construction"
FT	397. .532
FT	/note= "region of fusion derived from eta chain, preferred signal-transducing portion for constructs of the invention are amino acids 421-532, 423-455, 438-555, 461-494, 494-528 or 400-420;"
FT	Domain
FT	400. .437
FT	/label= Transmembrane domain
FT	/note= "eta chain transmembrane domain"
FT	438. .575
FT	Domain
FT	/label= Intracellular domain
FT	/note= "eta chain intracellular domain"

XX WO9625953-A1.  
 XX PD 29-AUG-1996.

PP 25-JAN-1996; 96WO-US001056.  
 XX DR 24-FEB-1995; 95US-00394176.  
 PR XX (GEHO ) GEN HOSPITAL CORP.  
 PA XX PI Seed B, Romeo C, Kolanus W;  
 XX DR WPI; 1996-402134/40.  
 DR XX N-PSDB; AAT56760.  
 PT Direction of cellular immune response using therapeutic cell expressing 2  
 chimeraic receptors - comprising region binding to target cell and region  
 that signals target cell destruction, or CD28 region, partic. for  
 PT eliminating HIV-infected cells.  
 XX  
 PS Claim 7; Page 77-78; 120pp; English.  
 XX  
 CC A chimaeric receptor (AAW00215) comprises the extracellular domain of an  
 engineered form of the CD4 cellular receptor for HIV and the  
 transmembrane and intracellular regions, including the cyrolytic signal-  
 transducing portion, of the mouse T-cell receptor eta chain. It can be  
 obtdd. by inserting a gene fusion (AAT36760) into a vaccinia vector  
 and expression in host cells. Chimaeric receptors comprising CD4 fused to  
 eta, eta (see also AAW02213) or Fc receptor gamma (see also AAW02214)  
 chains are capable of directing cytotoxic T lymphocytes to specifically  
 recognise and kill cells expressing HIV gp120, thus providing a therapy  
 for AIDS. (Updated on 16-OCT-2003 to standardise OS field)  
 CC  
 XX SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;  
 Best Local Similarity 99.1%; Pred. No. 9.2e-58;  
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy	2 RRSRSARPAPVQGQVOLYNLTNGREEVYDVLKRGGRGROPEMGGPKRRKNPORGLYNEL	61
Ds	423 KFSRSAFPVQOGONHOLYNLNLGRREEVDVLDRKGGRGROPEMGGPKRRKNPORGLYNEL	482
Qy	62 QDKXMAEYSEIGMKERRRKGHDQIYQGLSTATKTDIDALHQIQLPPR 111	
Ds	483 QDKXMAEYSEIGMKERRRKGHDQIYQGLSTATKTDIDALHQIQLPPR 532	

RESULT 12  
 AAW83141  
 ID AAW83141 standard; protein; 532 AA.  
 XX AC AAW83141;  
 XX DT 03-FEB-1999 (first entry)  
 DE Chimeric receptor containing human eta polypeptide.  
 XX Human; zeta; eta; gamma; membrane-bound chimeric receptor; infection;  
 KW tumour; cancer cell; autoimmune-generated cell; T cell receptor; CD3;  
 KW CD4; B cell receptor; Fc receptor; pathogen; bacterial; fungal;  
 KW protozoan; viral.  
 XX OS Synthetic.  
 OS Homo sapiens.  
 XX PN US5843728-A.  
 XX PD 01-DEC-1998.  
 XX PF 05-APR-1995; 95US-00417495.  
 XX PR 07-MAR-1991; 91US-00655961.  
 PR 06-MAR-1992; 92US-00847566.  
 PR 28-FEB-1994; 94US-00203866.  
 XX PA (GEHO ) GEN HOSPITAL CORP.

XX  
 PT Romeo C, Kolanus W, Seed B;  
 XX WPI; 1999-04582/04.  
 DR N-PSDB; AAV70157.

PT Membrane-bound chimeric receptors - comprising extracellular portion  
 PT which recognises and binds a target cell and an intracellular portion of  
 PT e.g. a T-cell receptor.

XX  
 PS Claim 11; Col 45-48; 57pp; English.

XX  
 CC The present invention describes DNA encoding a membrane-bound chimeric  
 CC receptor comprising: (a) an extracellular portion that specifically  
 CC recognises and binds a target cell or a target infective agent; and (b)  
 CC an intracellular portion of a T-cell receptor CD3, zeta or eta  
 CC polypeptide, a B-cell receptor polypeptide or an FC receptor polypeptide.  
 CC The present sequence represents a chimeric receptor containing the human  
 CC eta polypeptide. Cells expressing chimeric receptors of the present  
 CC invention can be administered to mammals in order to destroy pathogens  
 CC (e.g. bacteria, fungi, protozoa or viruses, especially HIV), cancer cells  
 XX or autoimmune-generated cells

SQ Sequence 532 AA:

Query Match 98.6%; Score 583; DB 2; Length 532;  
 Best Local Similarity 99.1%; Pred. No. 9.2e-58; Mismatches 0; Indels 0; Gaps 0;  
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Ov 2 RRSRSABEPAYQOGONQLYNEILGRREYDVLKRRGDPGGKRRKNPQEGLNEL 61  
 :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:  
 Db 423 KRSRSAEPAYQOGONQLYNEILGRREYDVLKRRGDPGGKRRKNPQEGLNEL 62  
 |||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:  

Ov 62 QDKXMAEYSEIGMKGERRGKGHDGLYQGISTATKOTYDALHMQALPPR 111  
 |||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:  
 Db 483 QDKXMAEYSEIGMKGERRGKGHDGLYQGISTATKOTYDALHMQALPPR 112  
 |||||:|||||:|||||:|||||:|||||:|||||:|||||:  
 Sequence 112 AA:

XX  
 Query Match 96.8%; Score 572; DB 8; Length 112;  
 Best Local Similarity 97.3%; Pred. No. 2.4e-57; Mismatches 1; Indels 0; Gaps 0;  
 Matches 107; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Ov 2 RRSRSABEPAYQOGONQLYNEILGRREYDVLKRRGDPGGKRRKNPQEGLNEL 61  
 :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:  
 Db 3 KRSRSAEPAYQOGONQLYNEILGRREYDVLKRRGDPGGKRRKNPQEGLNEL 62  
 |||||:|||||:|||||:|||||:|||||:|||||:  

Ov 62 QDKXMAEYSEIGMKGERRGKGHDGLYQGISTATKOTYDALHMQALPPR 111  
 |||||:|||||:|||||:|||||:|||||:  
 Db 63 QDKXMAEYSEIGMKGERRGKGHDGLYQGISTATKOTYDALHMQALPPR 112  
 |||||:|||||:|||||:|||||:  
 Sequence 112 AA:

RESULT 13

ID ADL67239 standard; protein; 112 AA.

AC ADL67239;

XX DT 20-MAY-2004 (first entry)

XX DB Human CD3 zeta chain intracellular domain.

XX KW T cell receptor; TCR; CD3 zeta chain; co-stimulatory signalling region; binding element; immunostimulant; therapy; cancer; human.

XX OS Homo sapiens.

XX PN US2004043401-A1.

XX PD 04-MAR-2004.

XX PR 28-MAY-2003; 2003US-00448256.

XX PR 28-MAY-2002; 2002US-0383872P.

XX PA (SLOC ) SLOAN KETTERING INST CANCER RES.

XX PI Sadelain M, Brentjens R, Maher J;

XX DR WPI; 2004-225696/21.

XX PT New nucleic acid polymer encoding a chimeric T cell receptor having a zeta chain portion, useful for treating disorders where the immune response needs to be induced, such as cancer.

XX PS Disclosure; SEQ ID NO 12; 25pp; English.

XX  
 CC The invention relates to a nucleic acid polymer encoding a chimeric T cell receptor (TCR) which comprises human CD3 zeta chain intracellular domain, a co-stimulatory signalling region and a binding element that specifically interacts with a selected target. The methods and compositions of the invention are useful for treating disorders where the immune response needs to be induced, such as cancer. The present sequence is human protein related to the invention.

XX SQ Sequence 163 AA;

Query Match 96.8%; Score 572; DB 8; Length 163;  
 Best Local Similarity 97.3%; Pred. No. 3.8e-57; Mismatches 1; Indels 0; Gaps 0;  
 Matches 107; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY	2	RPSRSSEPPAYQQGQNQLYNEELNGRREYYDLKQRGRDPMEGKPRRNQEGLNEL	61
Db	3	:       :       :       :       :       :       :       :       :	
QY	54	KFSISADAPAXQGQSONOLYNLSNLGEREYYDLDKQRGRDPMEGKPRRNQEGLNEL	113
Db	62	OKDKRAEAYSIGMKCERRRGKGHDGLYQGLSTATKOTYDALHMQALPPR	111
114	OKDKRAEAYSIGMKCERRRGKGHDGLYQGLSTATKOTYDALHMQALPPR	163	

RESULT 15

CC associated with expression of a polypeptide in a test cell sample, a  
CC method for treating cancers and a method for inhibiting the expression of  
CC gene in a cell. The composition and methods are useful for detecting,  
CC diagnosing, preventing and treating cancers, especially lymphoma and  
CC leukemia. These may also be used in screening for agents that modulate  
CC cancer. The present sequence is a human CAP protein sequence. Note: The  
CC sequence data for this patent did not form part of the printed  
CC specification, but was obtained in electronic format directly from WIPO  
CC at [ftp://wipo.int/pub/published\\_pct\\_sequences](ftp://wipo.int/pub/published_pct_sequences)

ABO84748 standard; protein; 163 AA.  
ABO84748;  
18-NOV-2004 (first entry)  
Human cancer-associated protein HP23-013.2.  
Human; cancer-associated protein; cytostatic; cancer; leukaemia;  
lymphoma; CAP.

Search completed: March 7, 2005, 07:13:07  
Job time : 48.7139 secs

PF 17-FEB-2004; 2004WO-US004730.  
XX PR  
PR 14-FEB-2003; 2003US-00367094.  
PR 14-MAR-2003; 2003US-00388938.  
PR 15-APR-2003; 2003US-00417375.  
PR 13-JUN-2003; 2003US-00461862.  
PR 15-SEP-2003; 2003US-00663431.  
PR 15-DBC-2003; 2003US-00737318.  
XX PA  
PA (SAGR-) SAGRES DISCOVERY INC.  
XX PI Morris DW, Morris DW, Malandro MS;  
XX DR WPI; 2004-652914/63.  
DR N-PSDB; ABD33068.  
XX PT New isolated cancer-associated polynucleotides and polypeptides useful  
PT for diagnosing, preventing or treating cancers, especially lymphoma and  
PT leukemia, or in screening for agents that modulate cancer.  
XX PS claim 18; seqid 882; 310pp; English.

contiguous nucleotides of any of the 233 polynucleotide sequences given in the specification, or its complement. The nucleic acids encode cancer associated proteins. Also included are an expression vector comprising the isolated nucleic acid cited above, a host cell comprising the above recombinant nucleic acid or expression vector, a microarray for detecting a cancer-associated (CA) nucleic acid comprising at least one probe comprising at least 10 contiguous nucleotides of any of the above-mentioned nucleotide sequences, an isolated polypeptide (encoded within an open reading frame of a CA sequence selected from any of the 95 polynucleotide sequences as mentioned in the specification, or its complement), an isolated antibody, (or its antigen binding fragment) that binds to the above polypeptide, a hybridoma that produces the above monoclonal antibody, a pharmaceutical composition comprising the above antibody and a pharmaceutical excipient, a kit for detecting cancer cells (comprising the antibody cited above, methods for diagnosing cancer or for detecting the presence or absence of cancer cells in an individual, a method for inhibiting growth of cancer cells in an individual, a method for delivering a therapeutic agent to cancer cells in an individual, an electronic library comprising the above polynucleotide or polypeptide (or their fragments), methods of screening for anticancer activity or for a bioactive agent capable of modulating the activity of a CA protein (CAP), methods for detecting cancer